

## **Amendments to the Specification**

Please amend the specification on page 1, after the title as follows:

### **CROSS-REFERENCE TO RELATED APPLICATIONS.**

Please amend the specification on page 1, after the previously added phrase "This application is a § 371 of International Application No. PCT/EP00/08994, filed September 14, 2000.":

### **BACKGROUND OF INVENTION.**

Please amend the specification on page 2, after line 12 (the second paragraph) insert:

### **DEFINITIONS**

Please amend the specification on page 2, after line 24 (sixth paragraph) insert:

### **SUMMARY OF THE INVENTION**

Please amend the specification on page 3, after line 25 (sixth paragraph) insert:

### **DETAILED DESCRIPTION**

The first paragraph on page 12 has been amended as follows:

#### **Example 3: Database searches**

Sequence analysis is performed using the GCG software (Wisconsin Package Version10.0, Genetics Computer Group (GCG), Madison, Wisconsin). For identity searches, GenEMBL, the Arabidopsis thaliana database (<http://genome-www3.stanford.edu>), Kazusa Aragidopsis opening site (KAOS[[;]]<http://zearth.kazusa.or.jp/arabi/>) and Swissprot are used. Peptide sequences are analysed by GeneQuiz (<http://celunba.ebi.ac.uk:8765/gqsrw/submit>) or Expasy (<http://www.expasy.ch/>).